

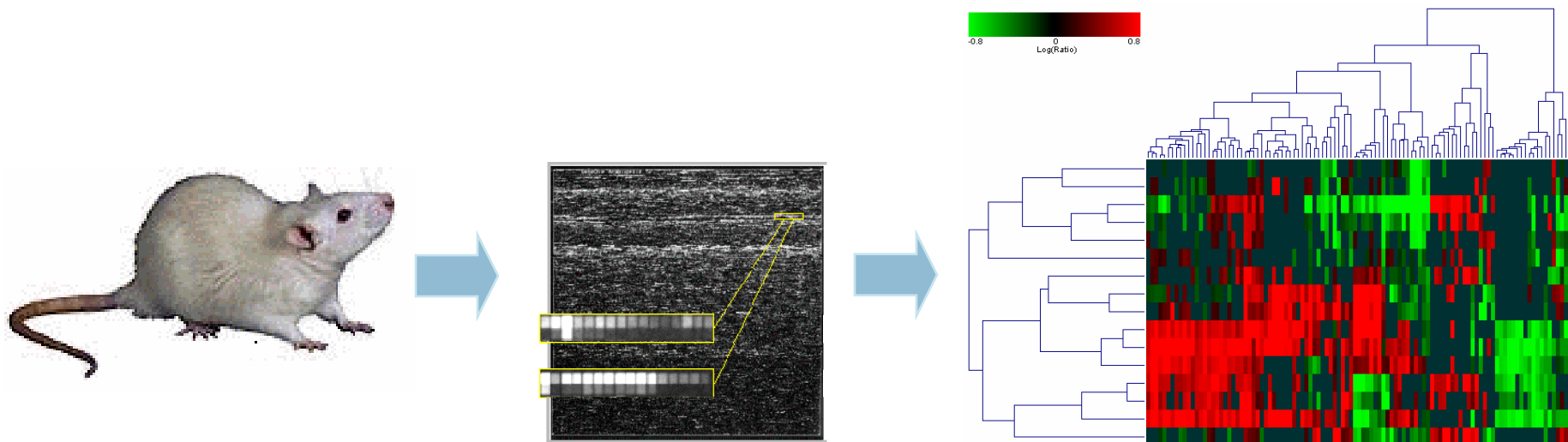


Application of *In Vitro* Toxicogenomics Towards Drug Safety Evaluation

Jeffrey F. Waring
Group Leader, Toxicogenomics
Abbott Laboratories

Toxicogenomics

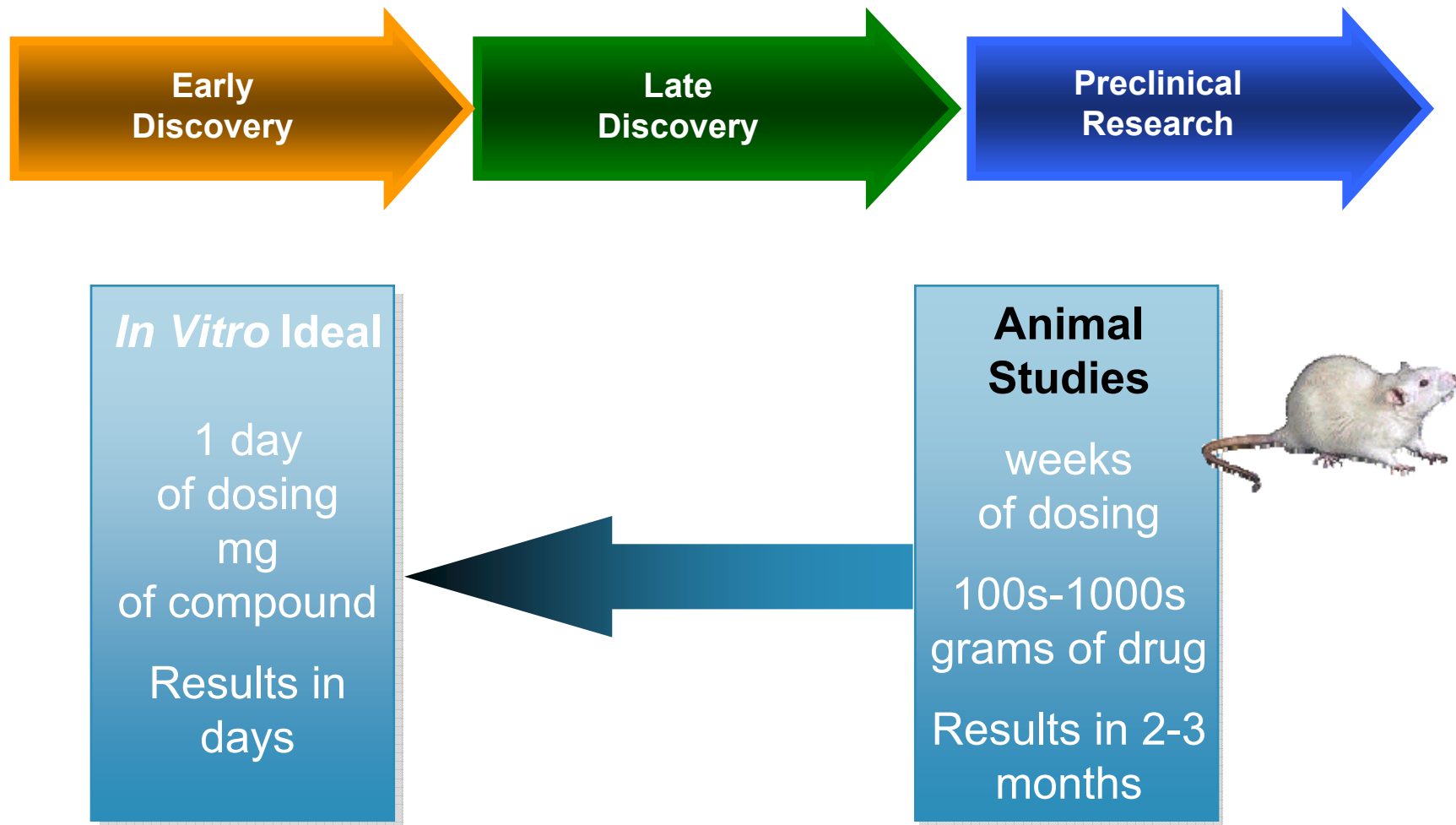
The application of gene expression analysis systems towards drug safety evaluation



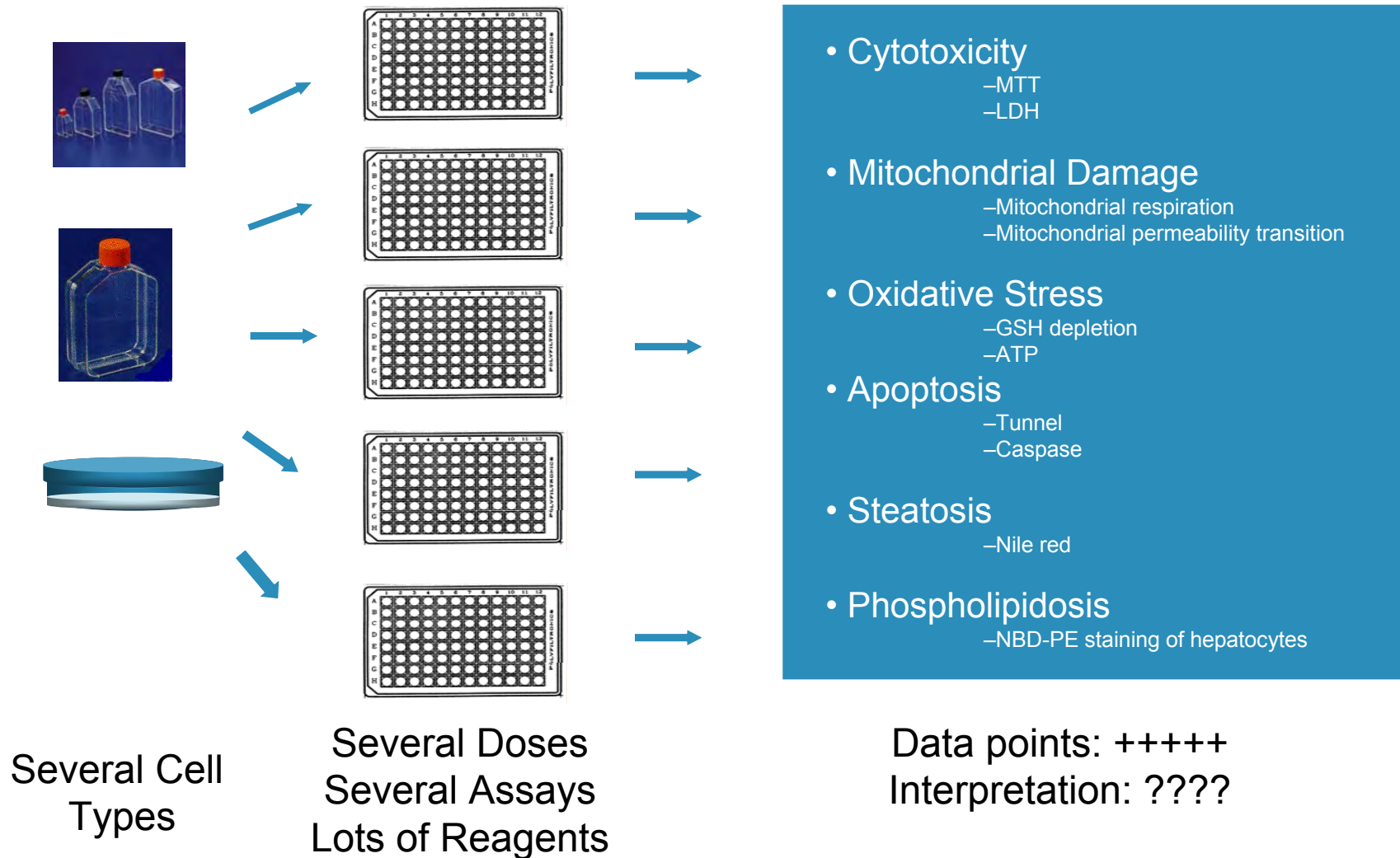
Information Gained from Toxicogenomics

- Patterns of gene expression changes associated with toxicity and with potential predictive value
- Specific gene expression changes related to the mechanism of toxicity
- Gene expression changes that can be used to bridge animal and human safety studies

Toxicological Characterization in Discovery

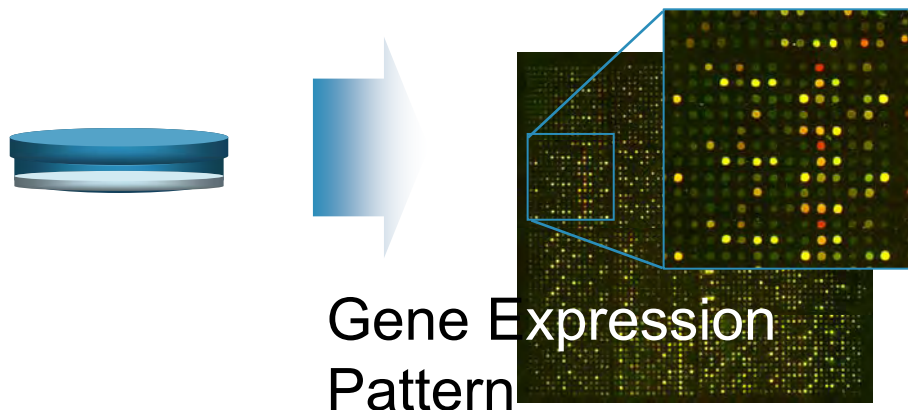


Traditional *In Vitro* Toxicology Paradigm



The *In Vitro* Toxicogenomics Paradigm

Collaboration with Iconix Pharmaceuticals



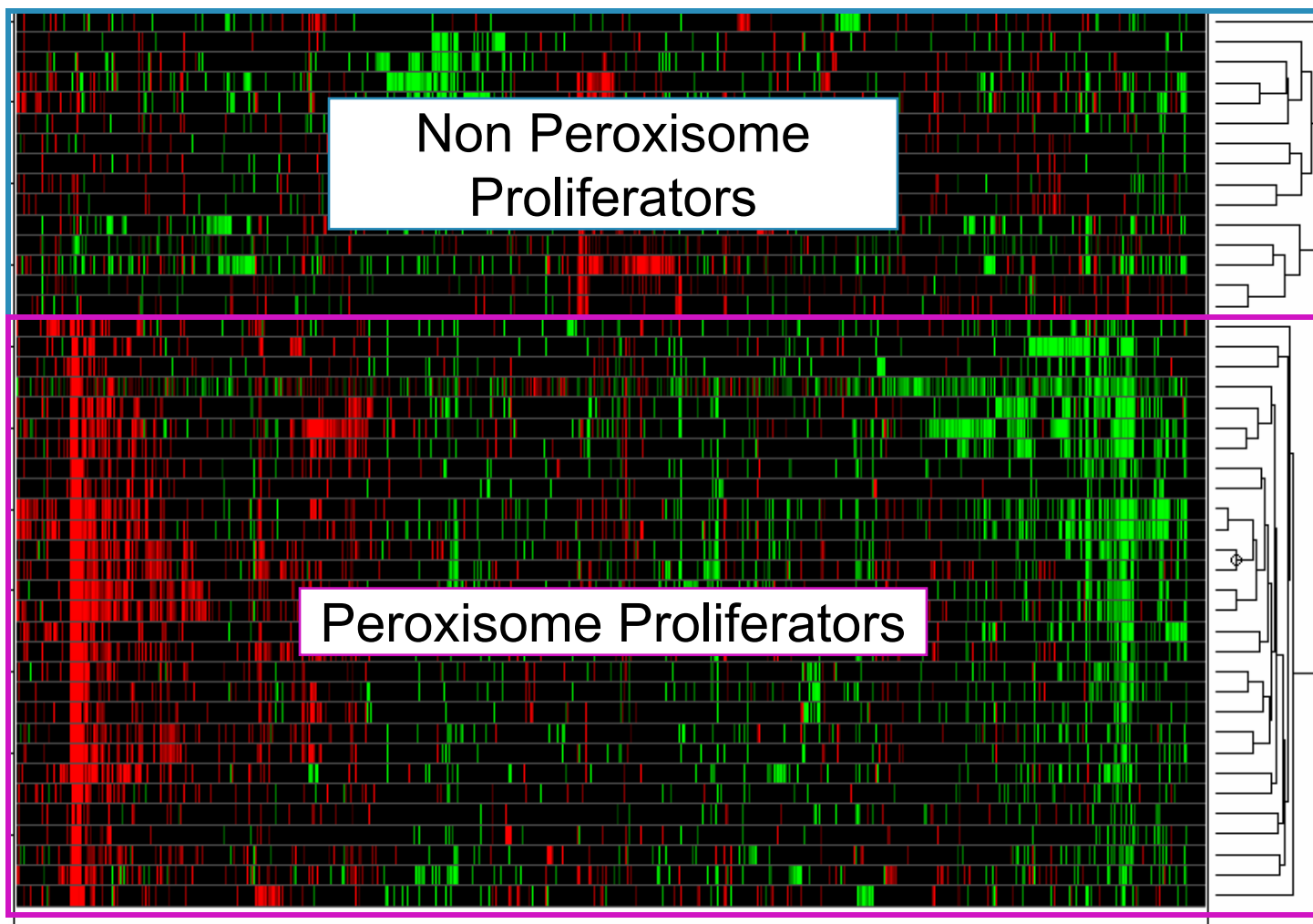
- Apoptosis
- Necrosis
- Canalicular cholestasis
- Microvesicular steatosis
- Peroxisome proliferation
- Ah-receptor agonist
- Phospholipidosis

One Cell
Type

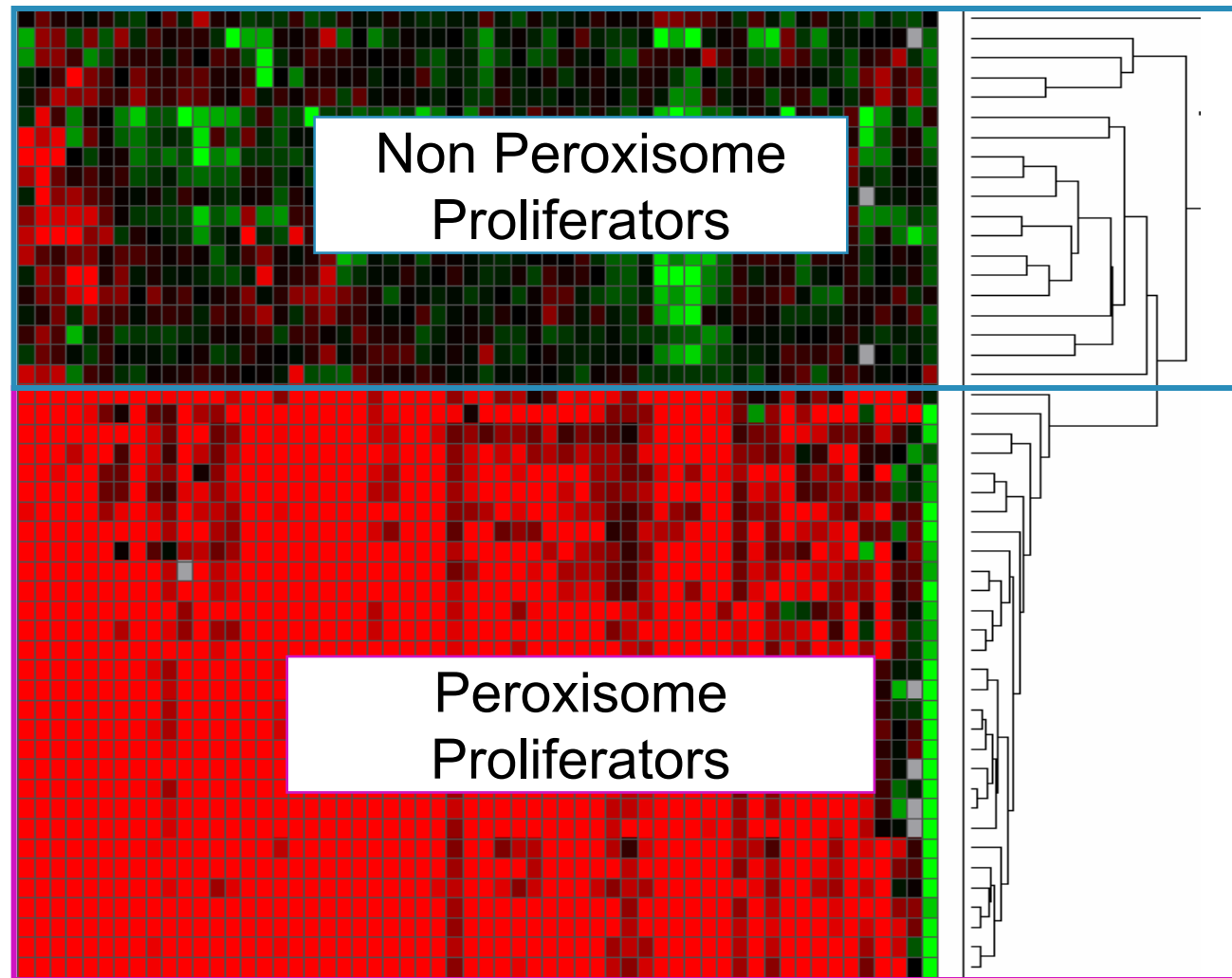
One Dose
One Assay
One Reagent Type

Data points: Limited
Interpretation: Simple

Mechanistically Similar Toxicants Induce Similar Gene Expression Changes *In Vivo*



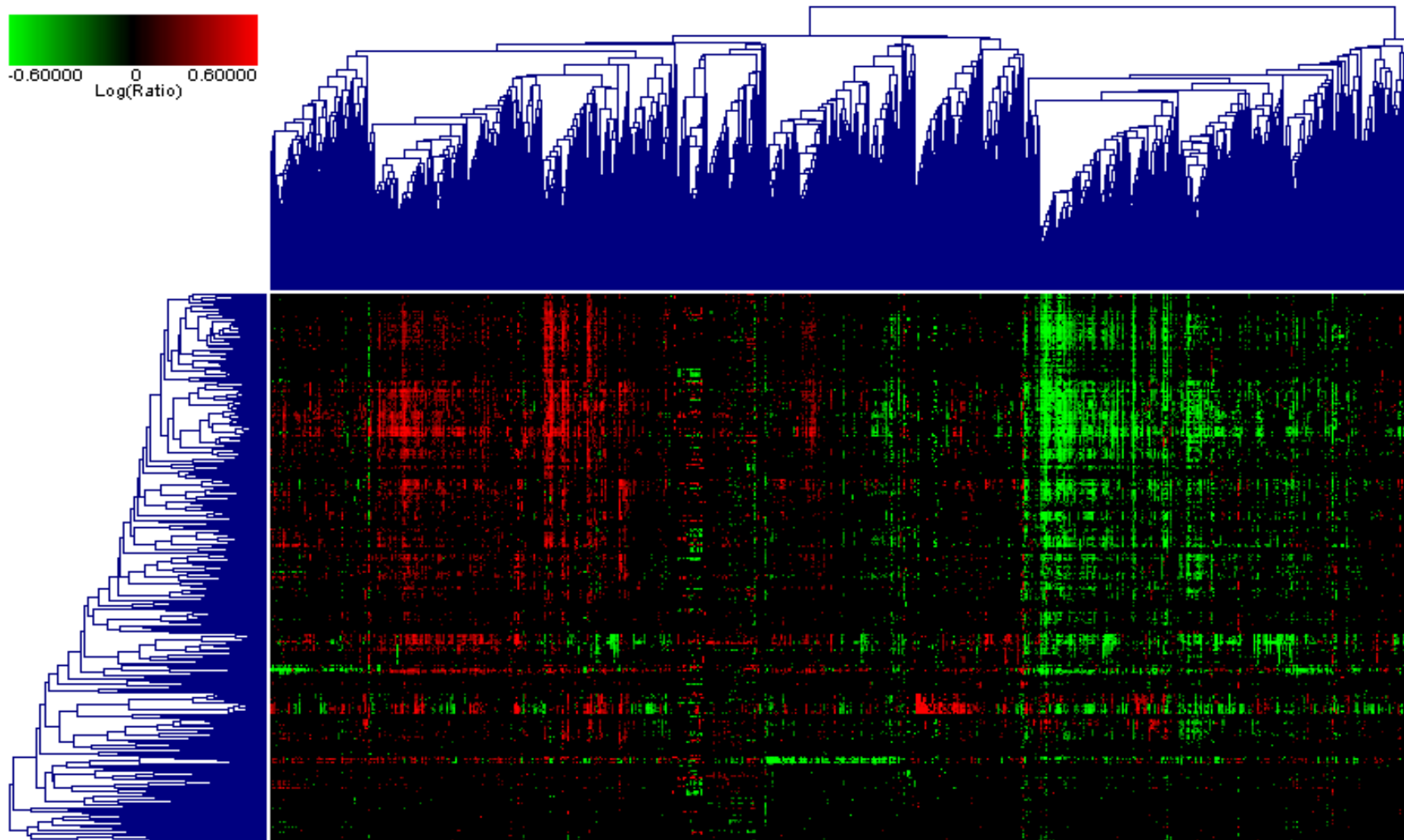
Signatures Can Be Generated for Mechanistic Class *In Vivo*



Toxicogenomics *in vitro* Assays: Rat Hepatocyte Protocol

- Isolated rat hepatocytes cultured for 24 hours before treatment
- Cells treated for 24 hours with compound at TC20 concentration
- 3 isolations used for all compounds
- Hepatocytes treated with compounds that are prototypical inducers of the toxicity
- Signatures created by identifying similar gene expression changes caused by compounds in the same mechanistic class
- 45 reference compounds
- 15 validation compounds
- 40 negative control compounds

Expression Profiles in Primary Rat Hepatocytes



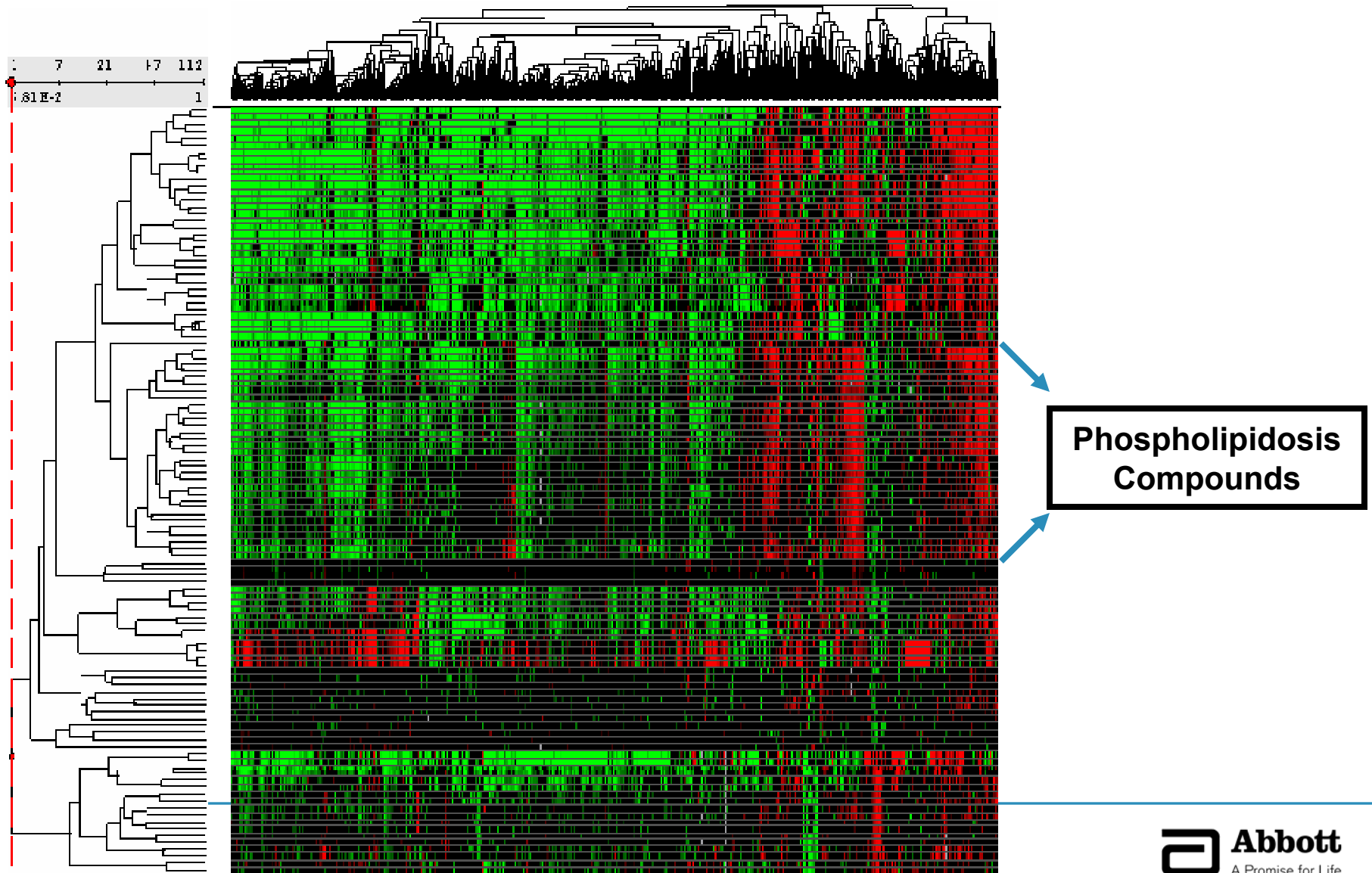
In Vitro Toxicogenomics

- Do compounds with similar mechanisms of toxicity give similar expression profiles *in vitro*?
- Can gene sets or signatures be identified that can be used to screen compounds?
- Can signatures be used to classify new compounds?
- What concentration should be used to screen new compounds?
- Can toxicogenomics be performed using assays that are more high throughput?
- How should these data be used for compound selection in drug discovery?

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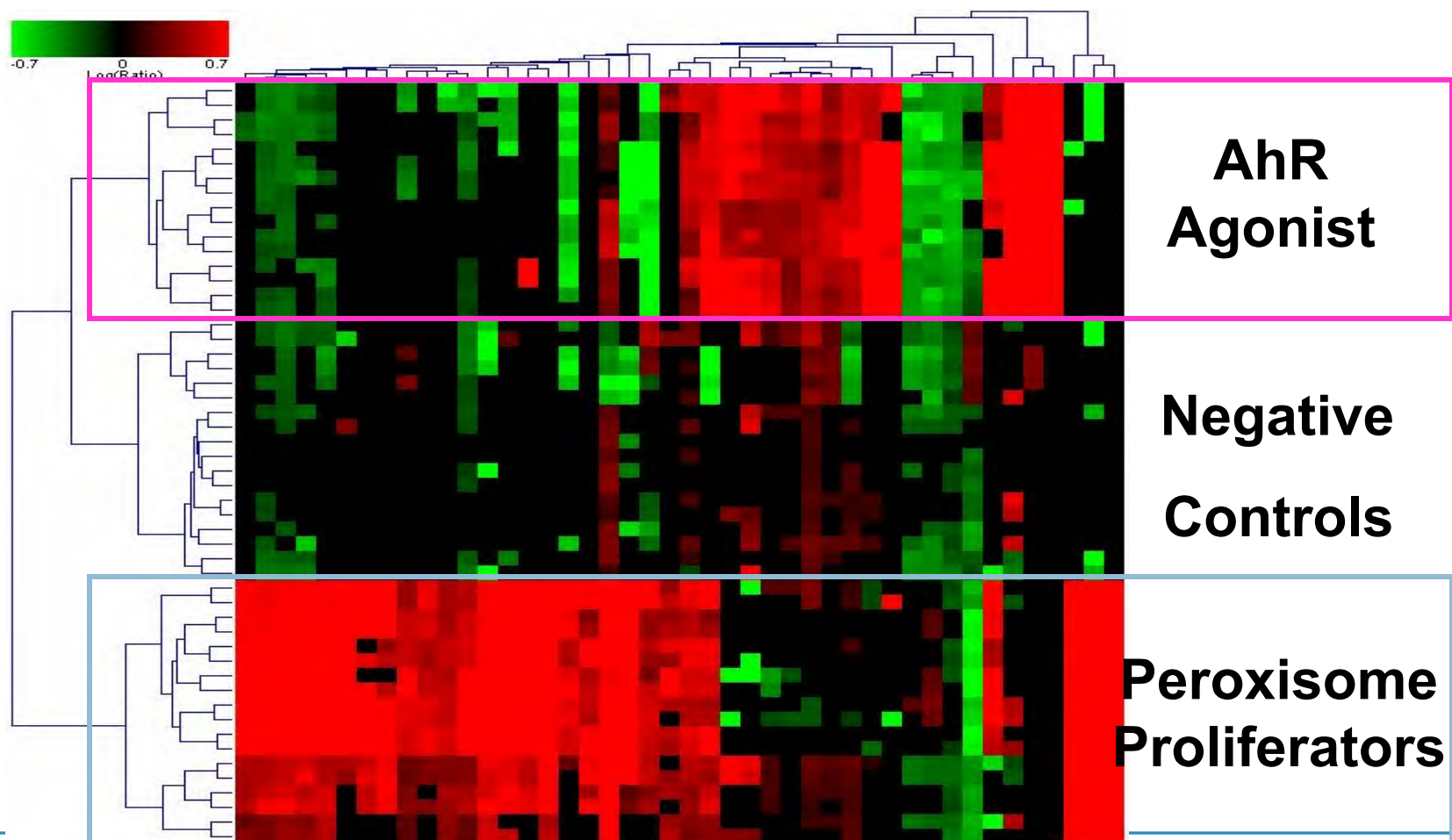
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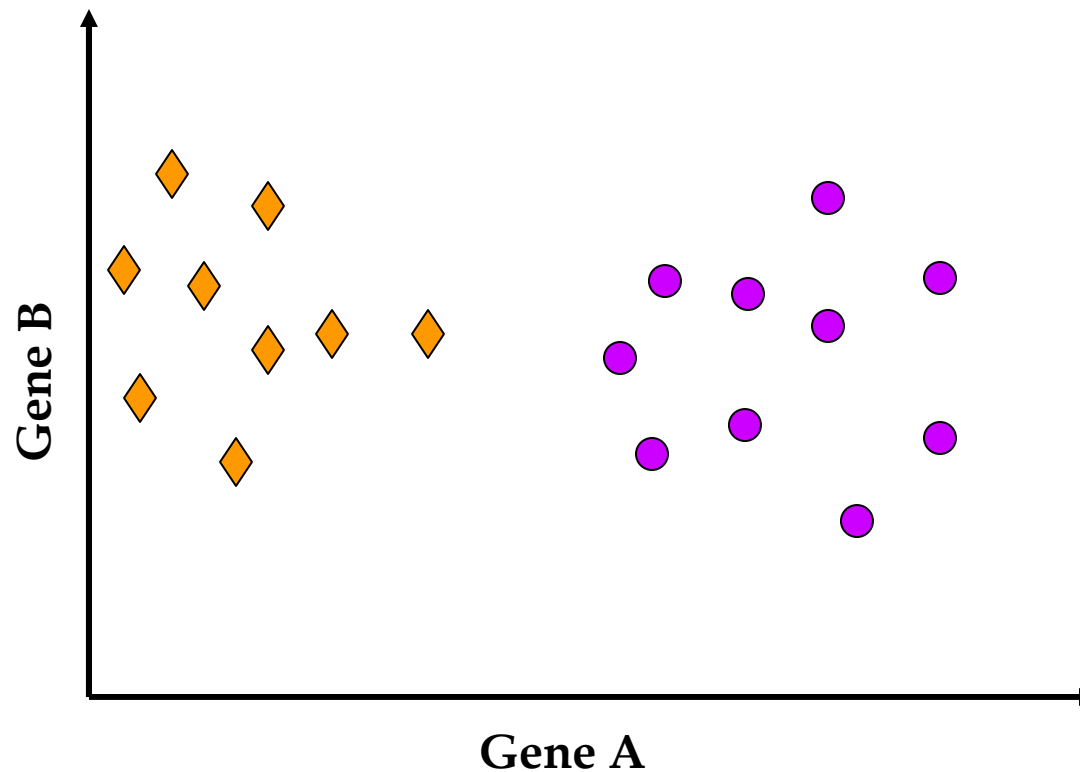
Compounds Tested for *In Vitro* Toxicogenomics

Compound Classes	Compounds
AhR agonist	3MC, Aroclor, Beta Naphthoflavone
Peroxisome Proliferator	Clofibrate, Bezafibrate, WY-14643
Negative Control	Penicillin, Spectinomycin, Chlorpheniramine

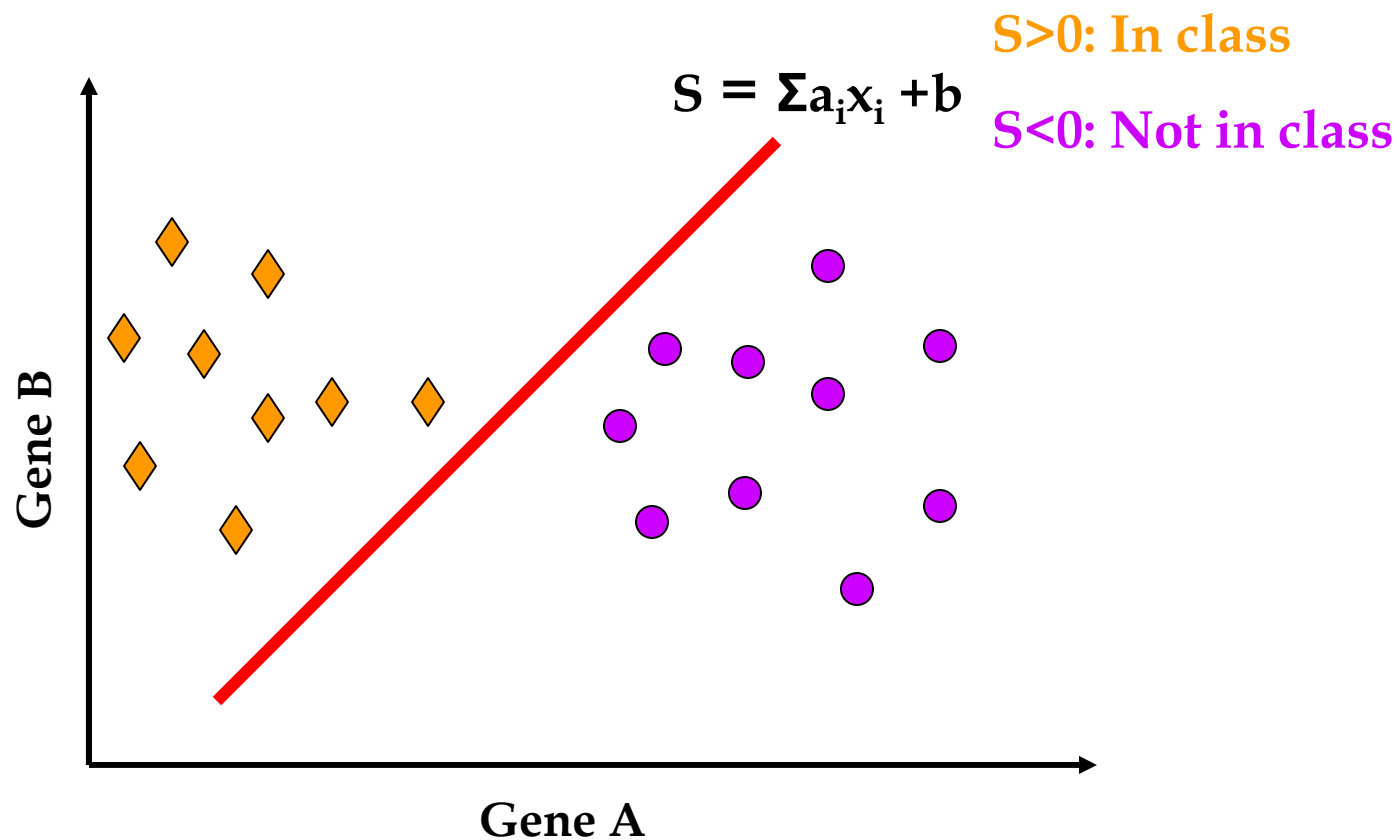
Compounds Classification using Hierarchical Clustering Analysis



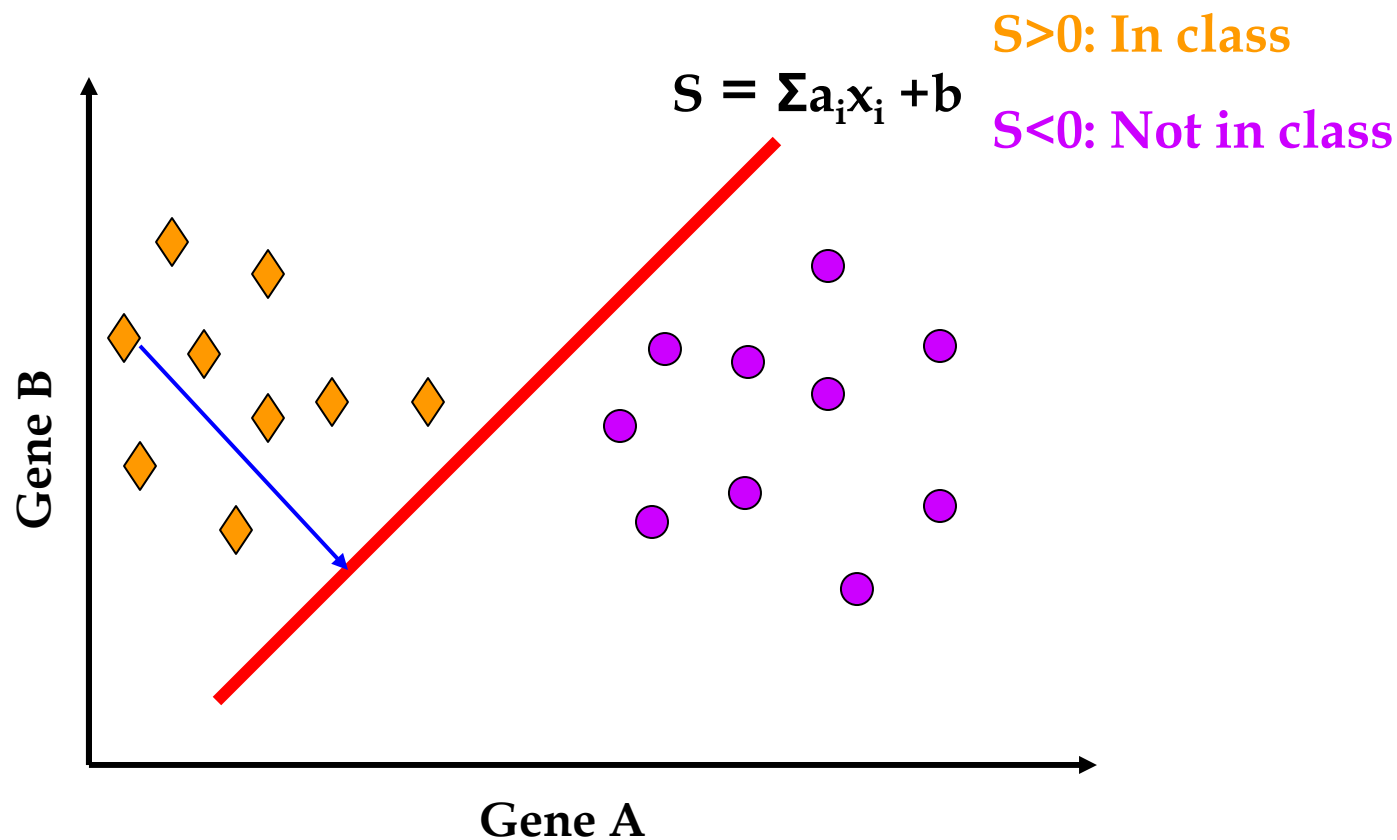
Linear Discriminant Analysis (LDA): a Statistical Algorithm for Classification



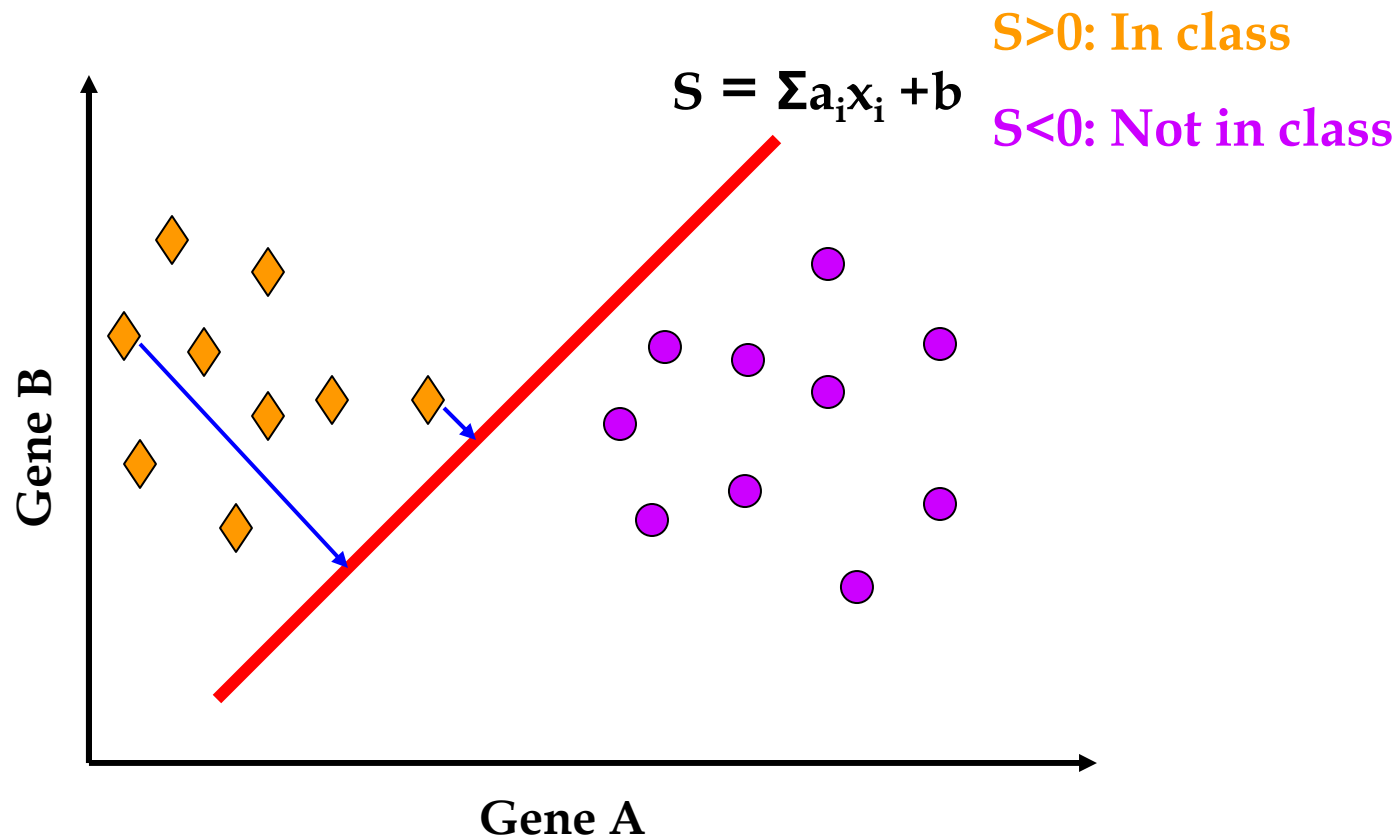
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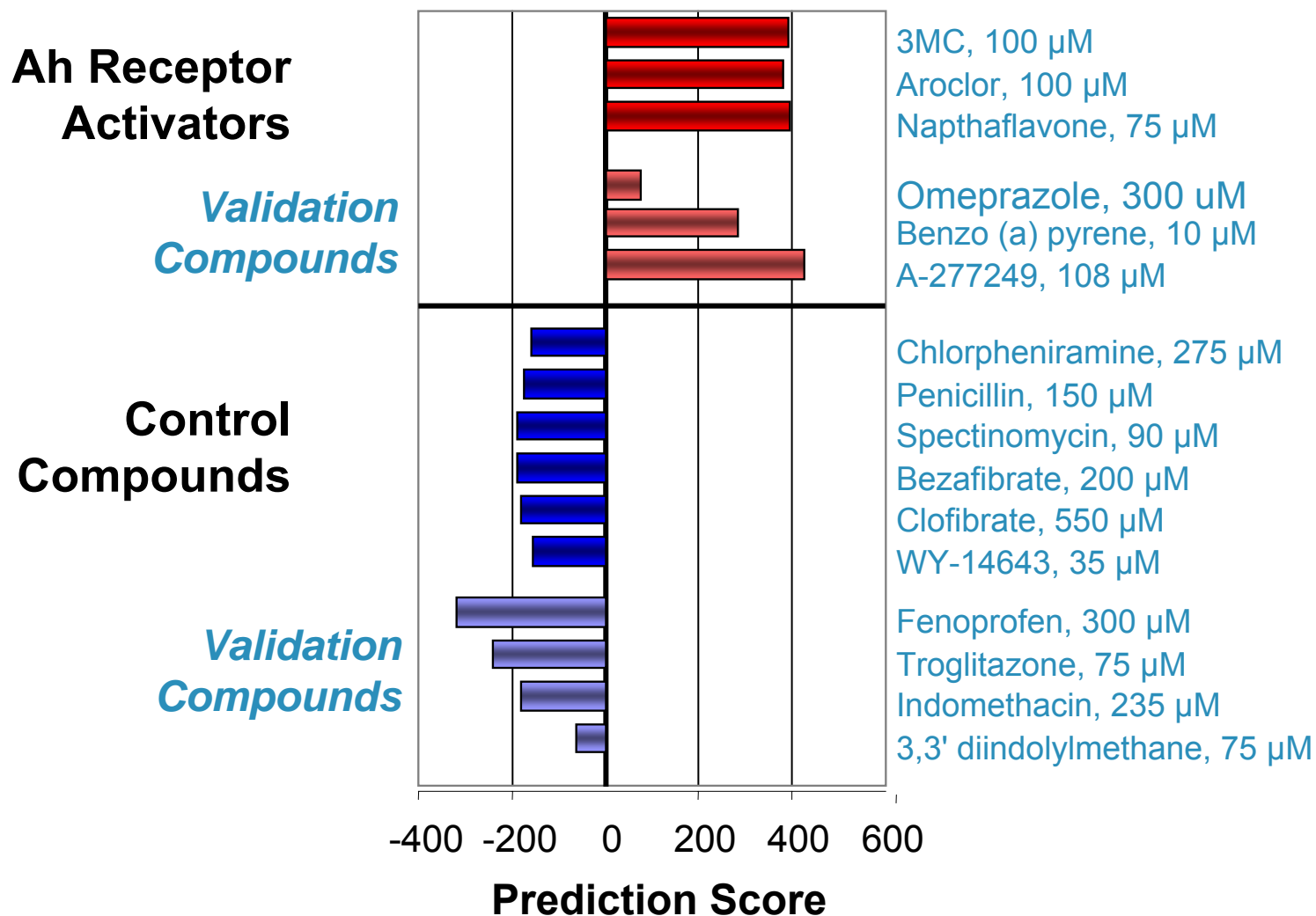
Validation Compounds for *In Vitro* Signatures

- **AhR Agonist**
 - Benzo(a)pyrene
 - A-277249
 - Omeprazole

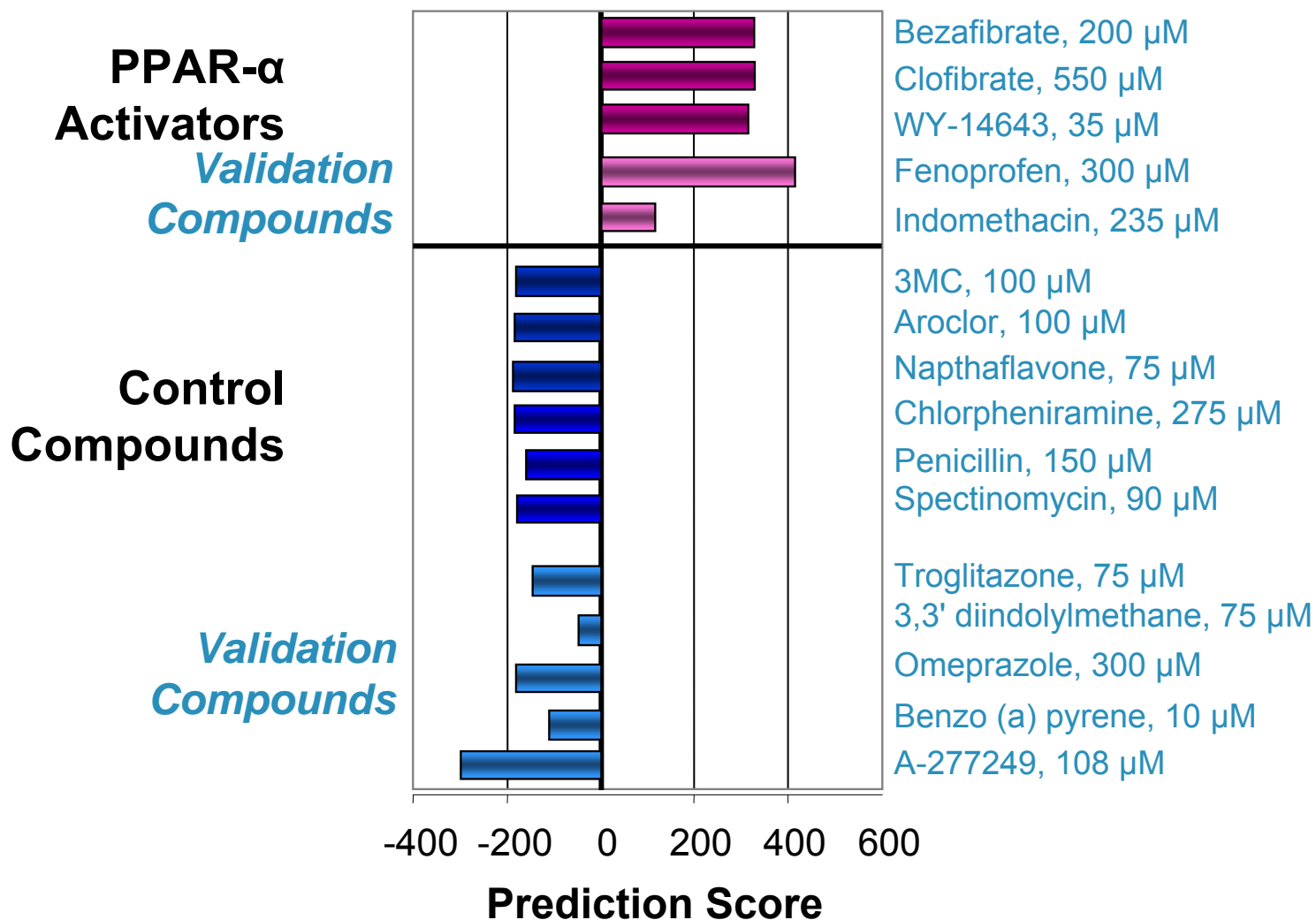
- **Peroxisome Proliferator**
 - Fenoprofen
 - Indomethacin

- **Negatives**
 - 3',3-diindolylmethane (DIM)
 - Troglitazone

In Vitro Signatures Correctly Classify Known Hepatotoxins



In Vitro Signatures Correctly Classify Known Hepatotoxins



In Vitro Toxicogenomics

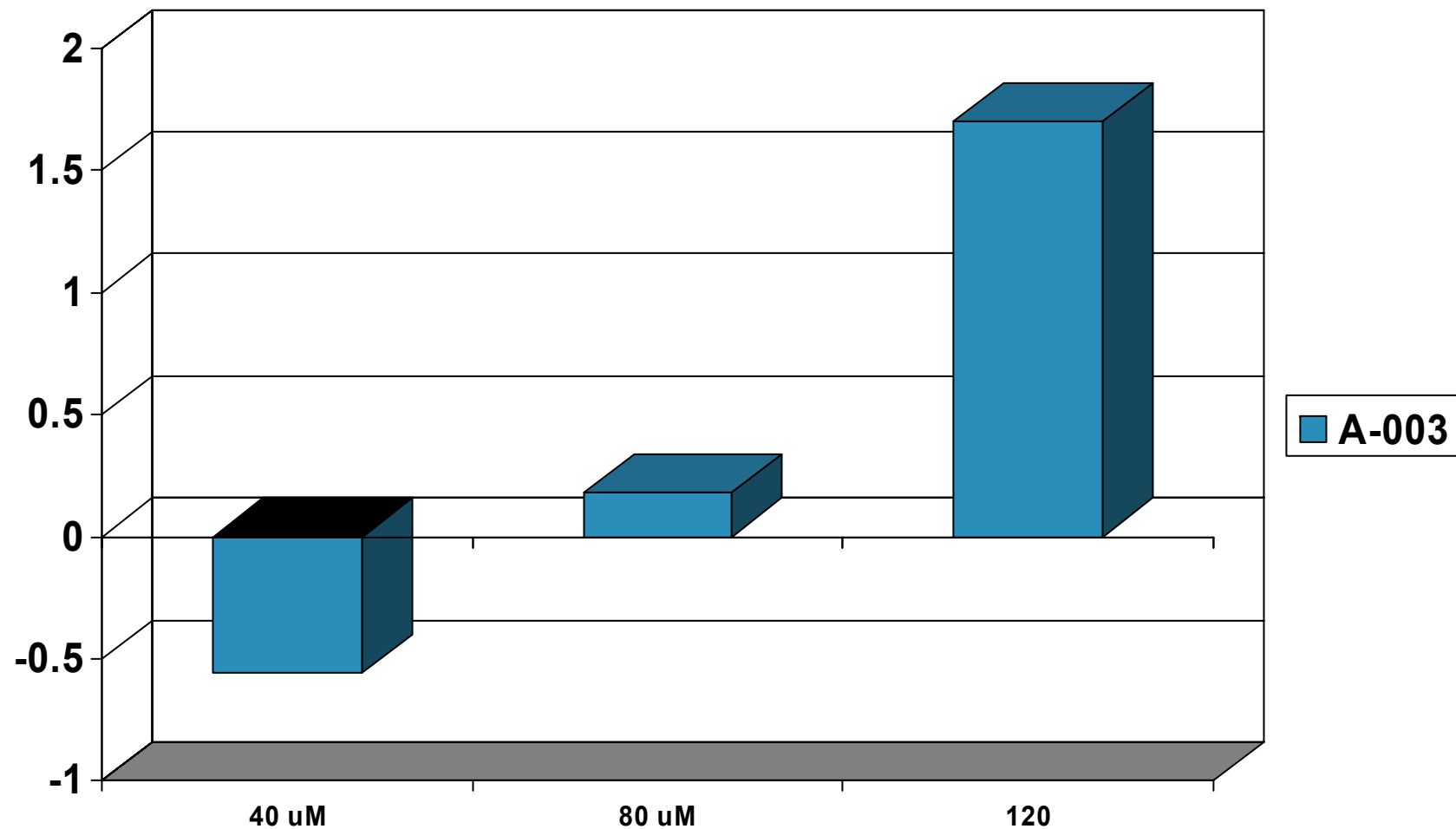
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Phospholipidosis Signature

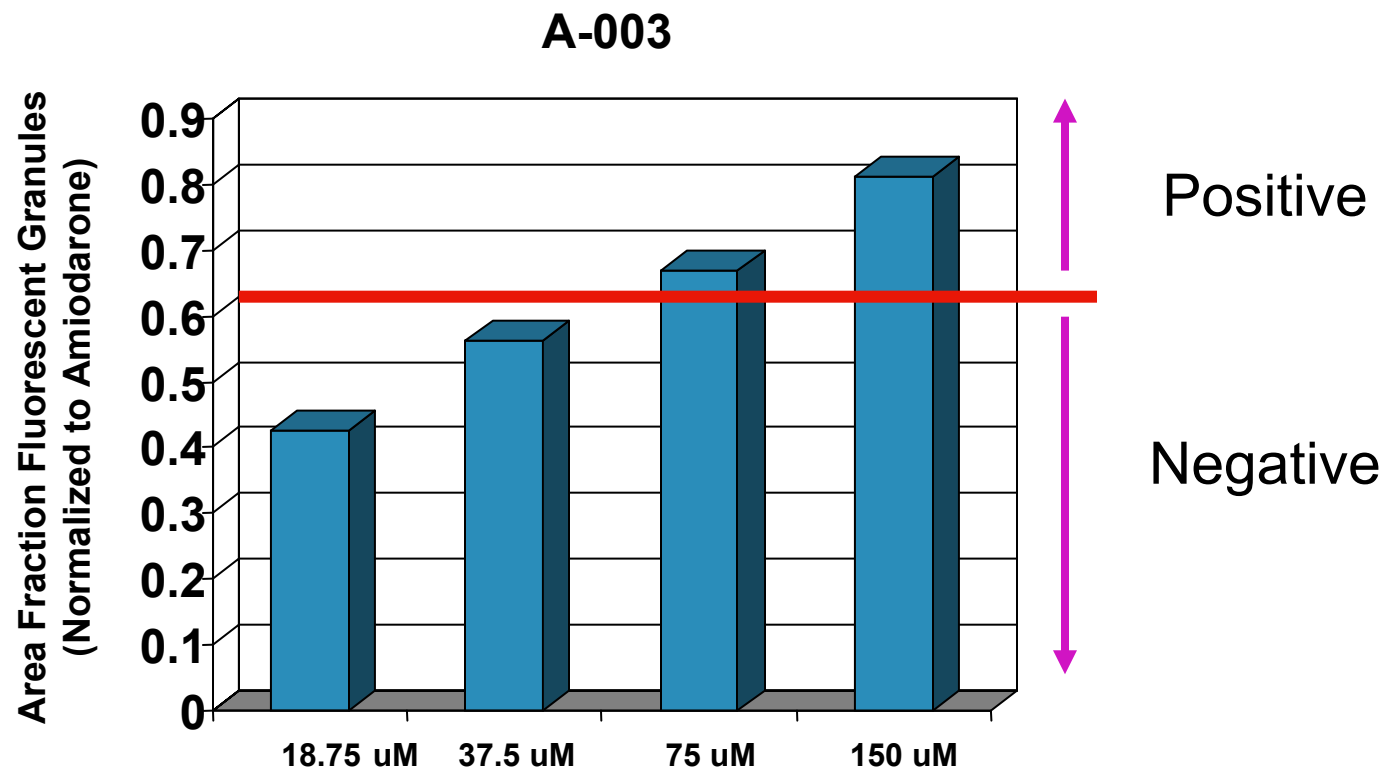
Compound	Dose	Result in Phospho. Cell Assay	Phospholipidosis Signature
A-001	40 μ M	+	+
A-002	40 μ M	++	++
A-003	40 μ M	+	-
Cyclophosphamide	1.32 mM [#]	NA	-
Doxorubicin	1.5 μ M [#]	NA	-
Methapyrilene	300 μ M [#]	NA	-
Rifampin	125 μ M [#]	NA	-

[#]: dose corresponding to TC20 at 24 hr.

Phospholipidosis Signature



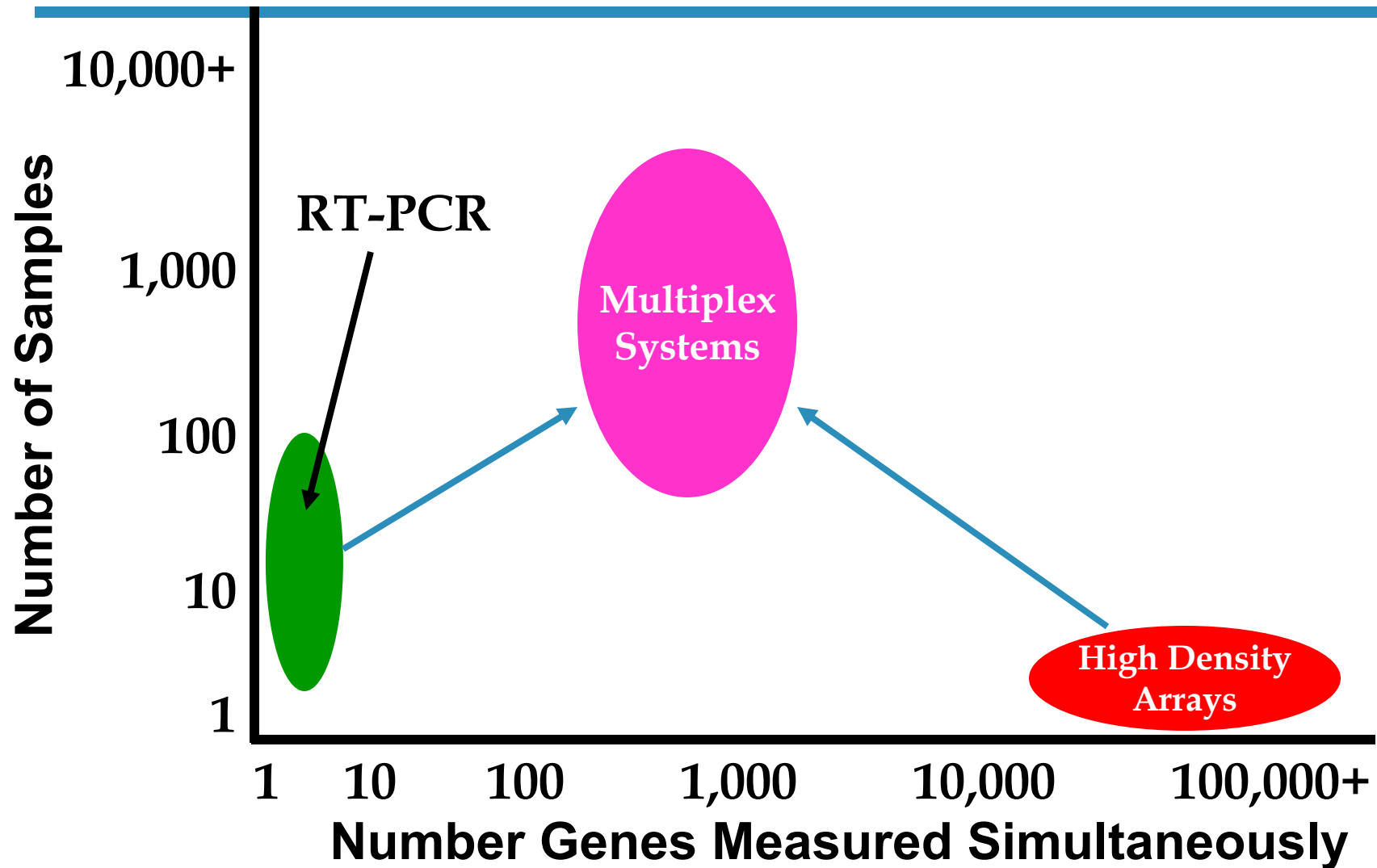
Phospholipidosis Cell Based Assay



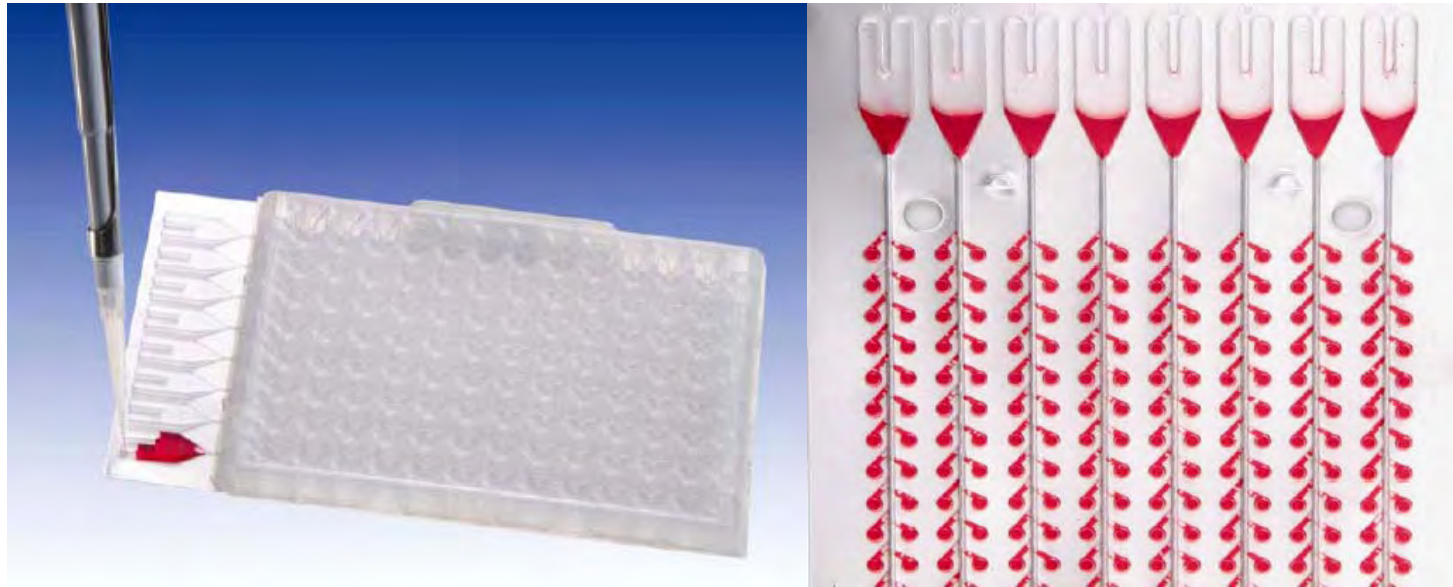
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Gene Expression Profiling: Moving Toward Higher Throughput

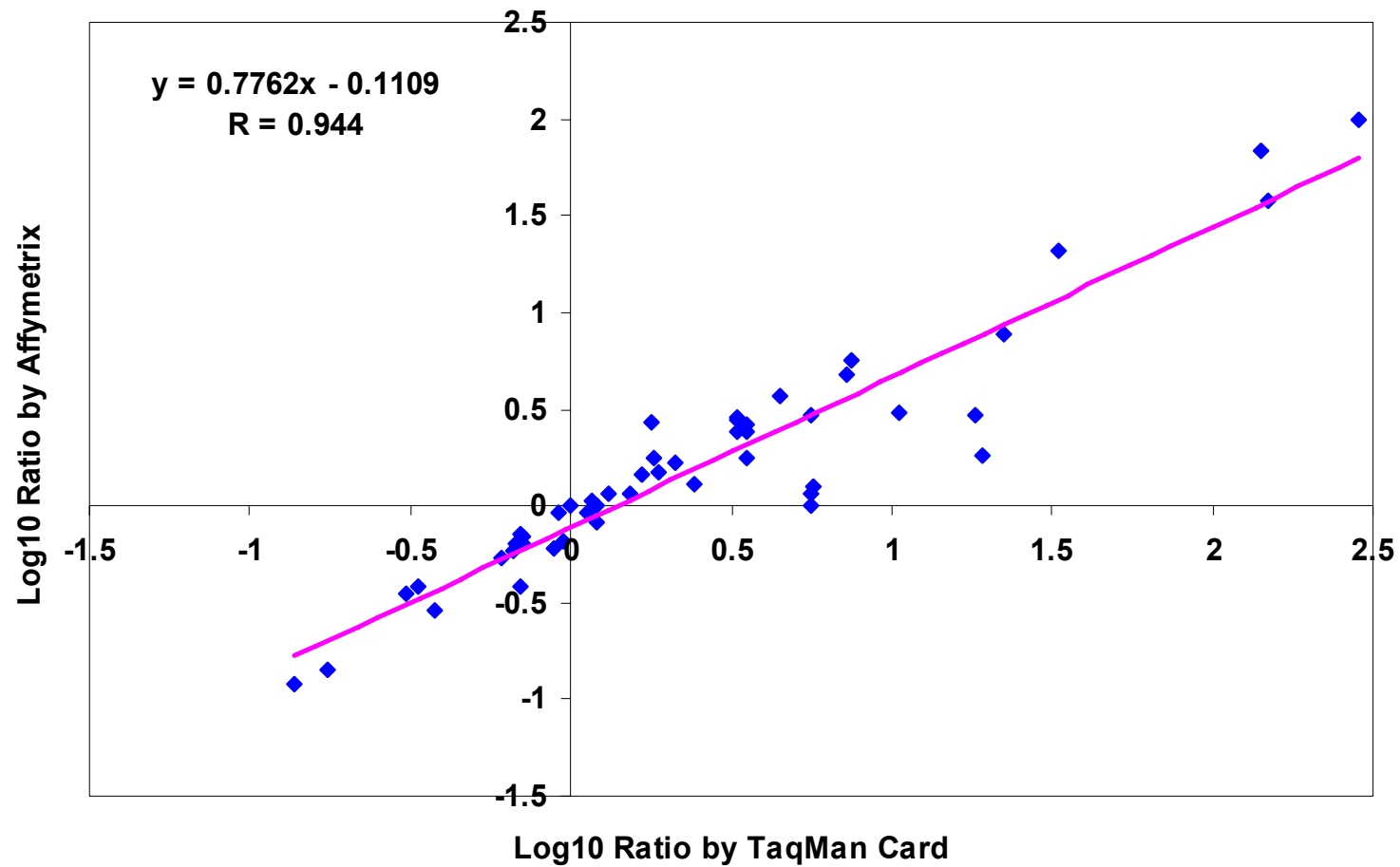


TaqMan Micro Fluidic Card



- Capable of identifying expression changes up to 200 genes
- Ability to process 20-50 samples in a week
- Cost under \$100 a sample
- Flexibility to add new genes

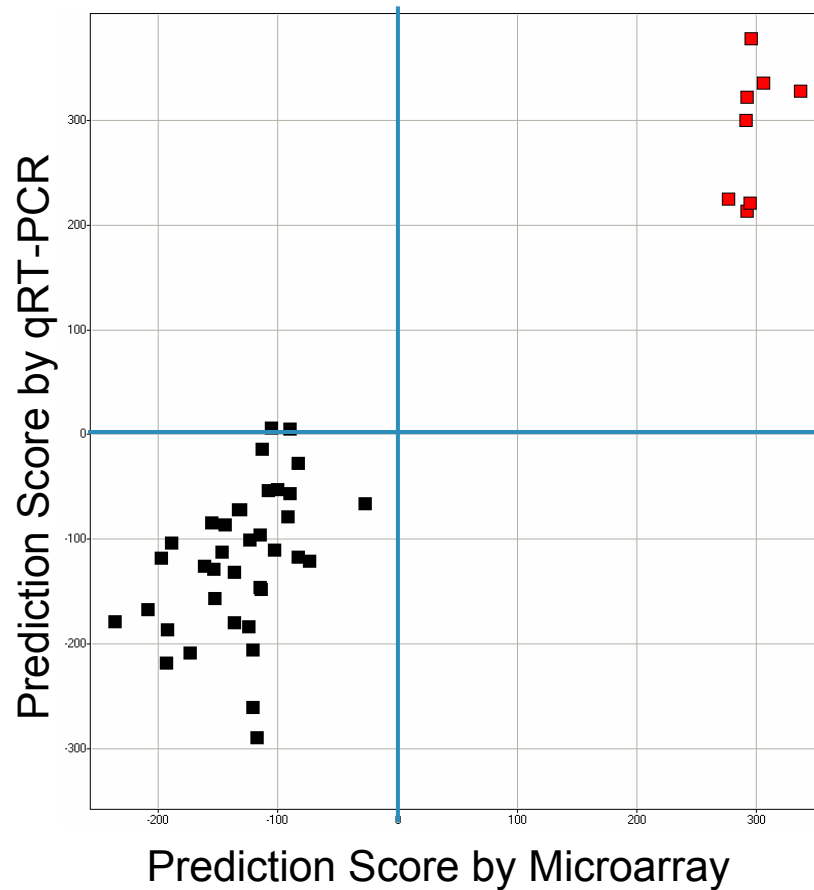
RT-PCR Card vs. Microarray



Prediction of AhR Activator



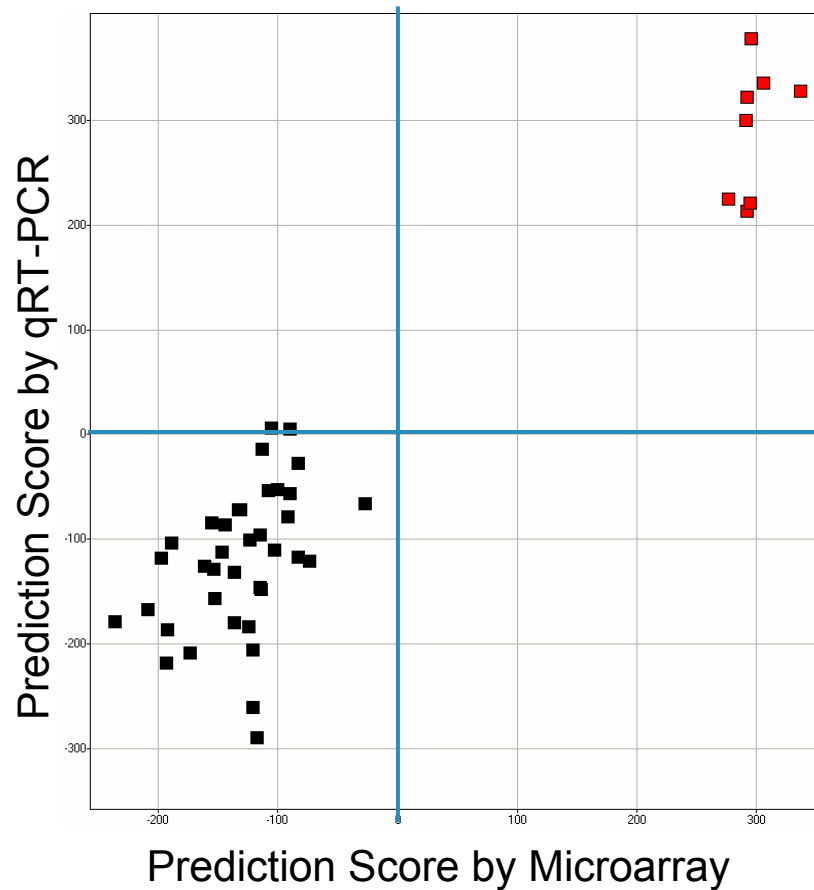
Training Set



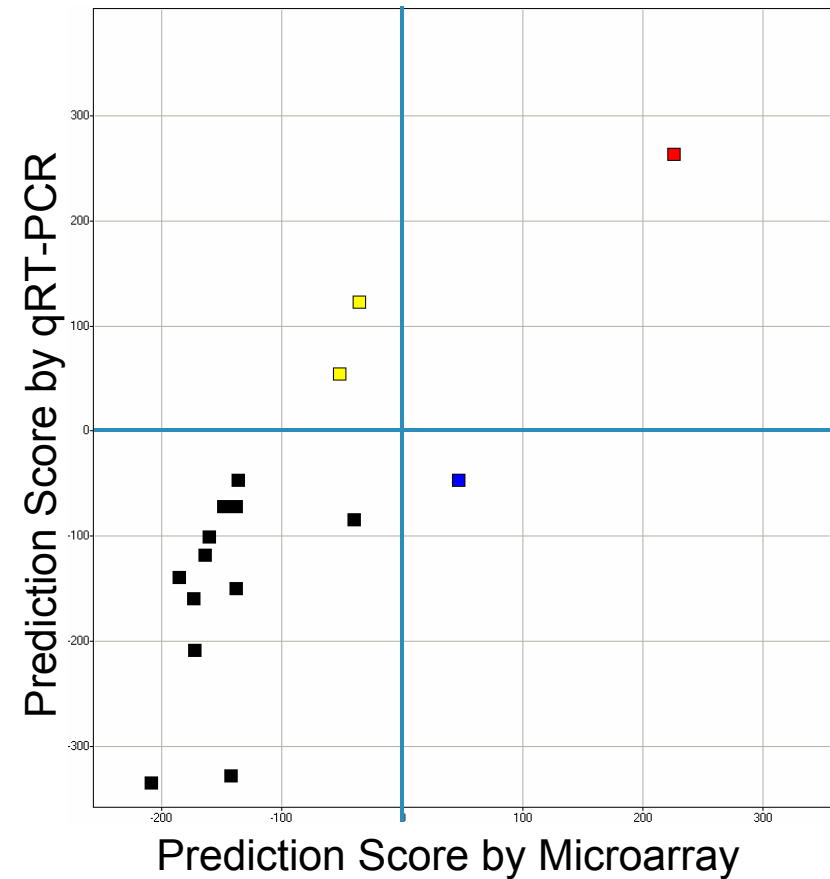
Prediction of AhR Activator



Training Set



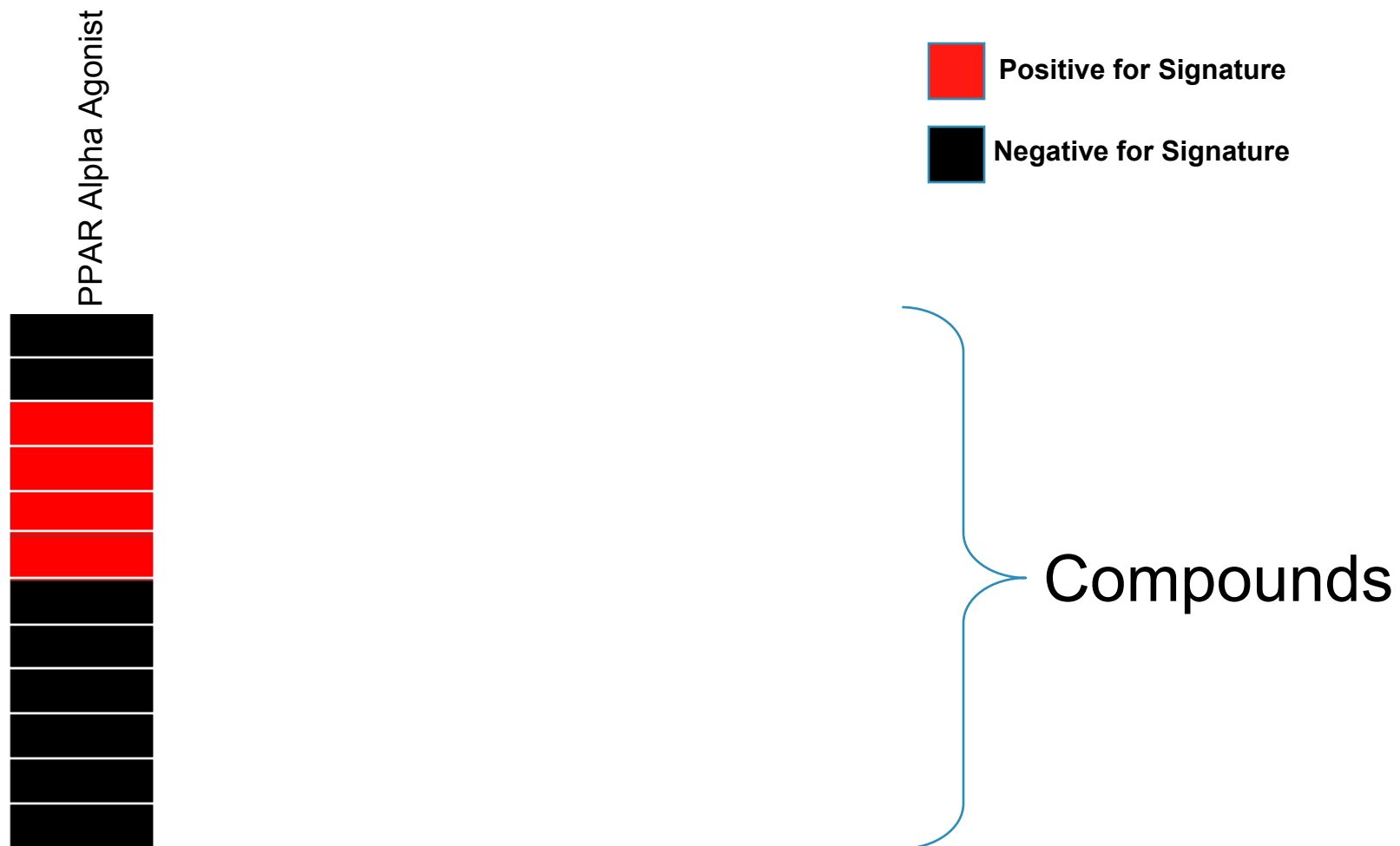
Validation Set



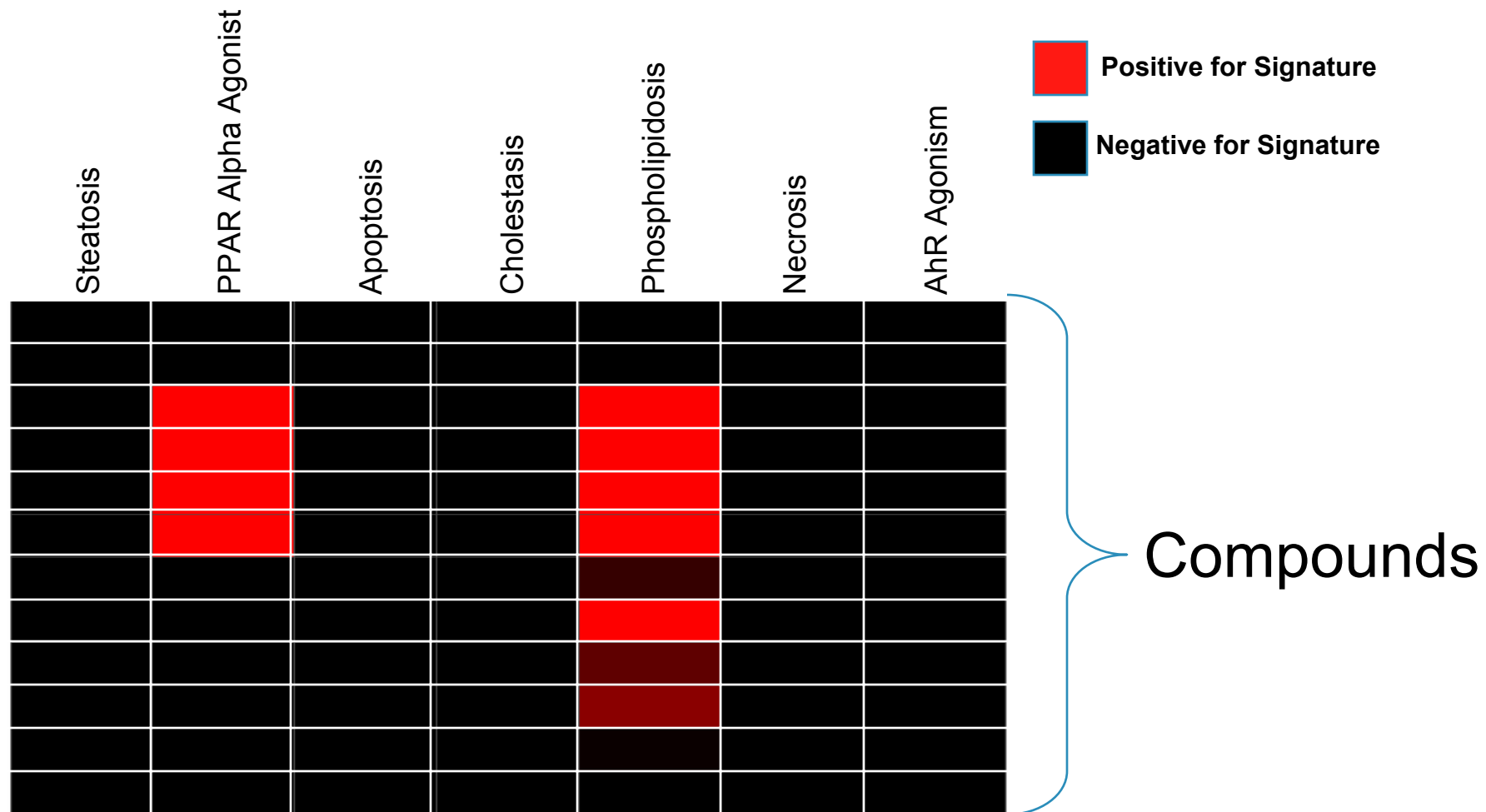
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Evaluation of Compounds Using *In Vitro* Toxicogenomics





Evaluation of Compounds Using *In Vitro* Toxicogenomics



Can Safety Margins Be Determined?

	Phospholipidosis	Steatosis	Perox. Prol.	Apoptosis	Cholestasis	AhR activation	Necrosis
Cpd 1 100 uM							
Cpd 1 50 uM							
Cpd 1 10 uM							

 Positive for Signature

 Negative for Signature

Human PBMCs for In Vitro Characterization

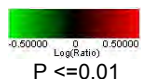
- Identify toxicities that may be more relevant for humans
- Human PBMCs would reflect genetic diversity present in human population
- Identify biomarkers that can be readily transferred to the clinic

In Vitro Screening Using Human PBMCs

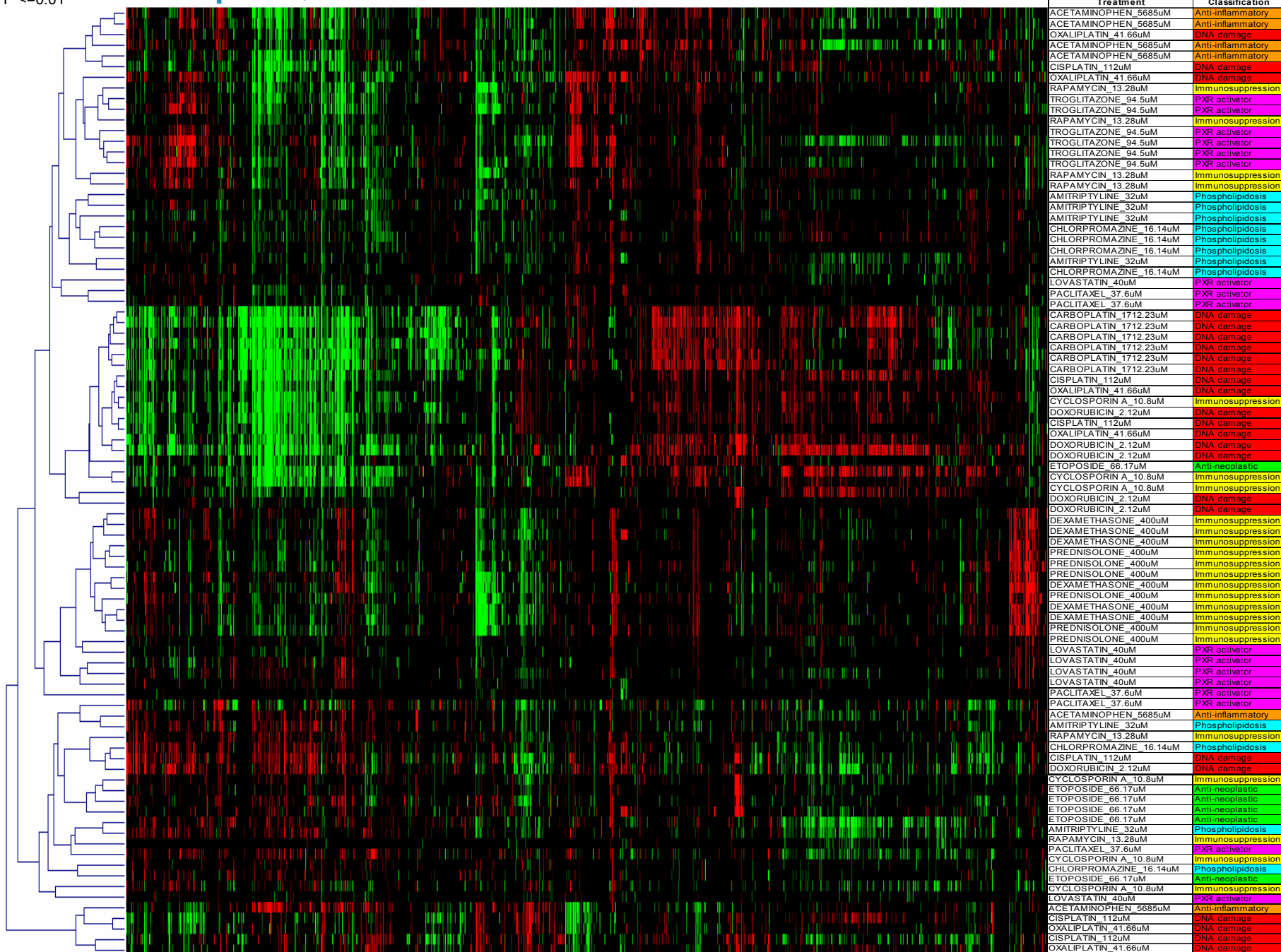
Compound_Name	Dose			Classification	Structure_Activity
	uM	MTD (TC20)	MFD		
DOXORUBICIN	3.59	Yes		DNA damage	DNA intercalator, anthracycline
CARBOPLATIN	1456.5	Yes			DNA-alkylator, platin
CISPLATIN	152.6	Yes			DNA-alkylator, platin
OXALIPLATIN	38.6	Yes			DNA-alkylator, platin
ETOPOSIDE	56.6	Yes		Anti-neoplastic	DNA topoisomerase II inhibitor
ACETAMINOPHEN	6509.2	Yes		Anti-inflammatory	NSAID, COX-3, acetaminophen like
PREDNISOLONE	400		Yes	Immunosuppression	Glucocorticoid and mineralocorticoid receptor agonist
CORTISONE	80		Yes		Glucocorticoid receptor agonist
DEXAMETHASONE	400	Yes	Yes		Glucocorticoid receptor agonist
CYCLOSPORIN A	8.58	Yes			Inhibits T-cell activation
CHLORPROMAZINE	25	Yes		Phospholipidosis	Dopamine receptor antagonist (D), phenothiazine
RIFAMPIN	80.25	Yes		PXR activator	RNA polymerase inhibitor
CLOTRIMAZOLE	17.6	Yes			Sterol 14-demethylase inhibitor
BENZO[A]PYRENE	80		Yes	AhR Agonist	Toxicant, Ah receptor agonist

In Vitro Screening Using Human PBMCs

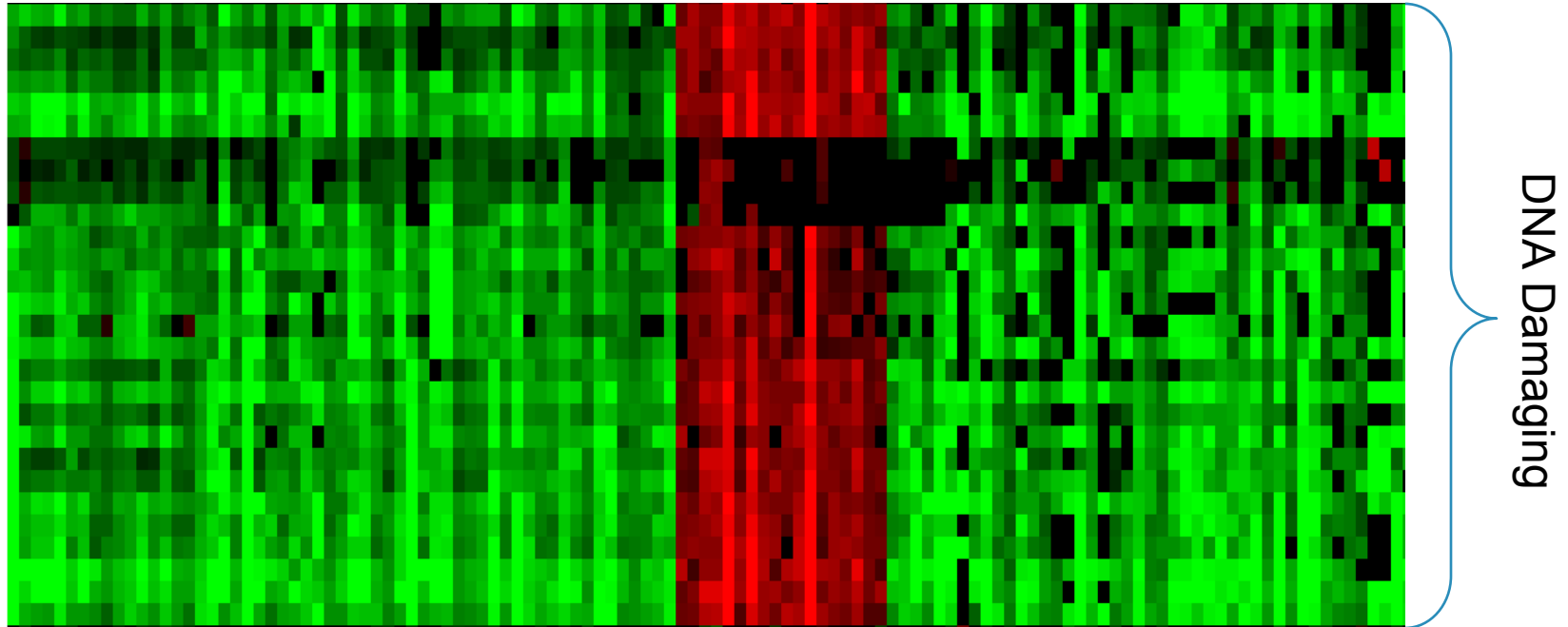
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PREDNISOLONE	400		Yes	Immunosuppression
CORTISONE	80		Yes	
DEXAMETHASONE	400	Yes	Yes	
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CHLORPROMAZINE	25	Yes		Phospholipidosis
RIFAMPIN	80.25	Yes		PXR activator
CLOTRIMAZOLE	17.6	Yes		
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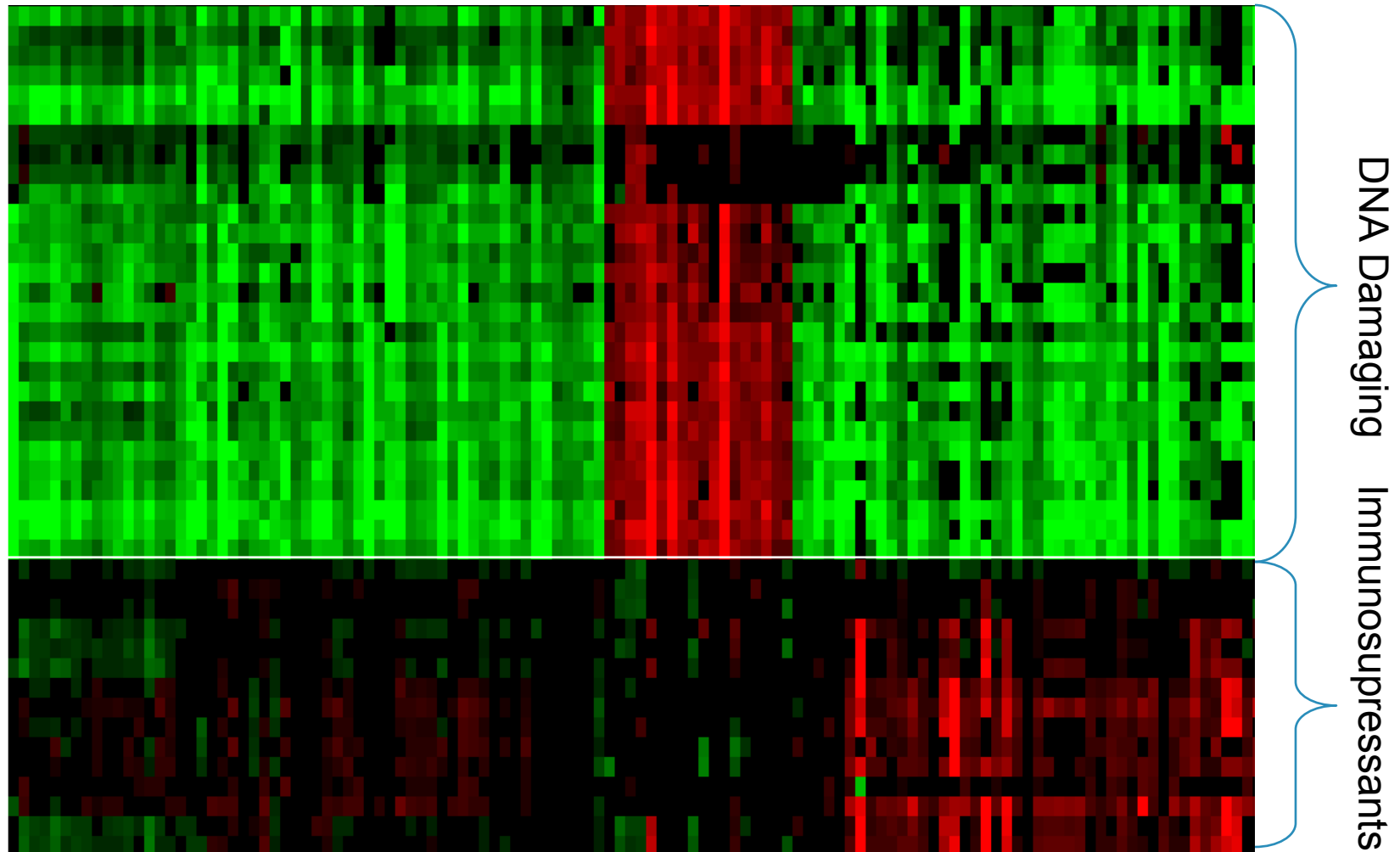
15 Cpds, 5~6 Donors (FC>=2 and p<=0.01 in at least one experiment, n=14,628)



DNA Damaging Agents Versus Immunosuppressants



DNA Damaging Agents Versus Immunosuppressants



Summary

1. In vitro toxicogenomics is a useful tool for SAR, prioritization of compounds, selection of backup compounds
2. Limitation is that safety margins in vivo cannot be determined
3. Together with other molecular and cell-based ADMET methods, these efforts should help shift attrition earlier in Drug Discovery

Acknowledgements

Cellular and Molecular Toxicology

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